



#9

SEQUENCE LISTING

<110> Bristol-Myers Squibb Company

<120> A NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPRBMY6, EXPRESSED HIGHLY IN SMALL INTESTINE

<130> D0040NP/3053-4119US3

<140> US 09/966,422

<141> 2001-09-26

<150> 60/235,602

<151> 2000-09-27

<150> 60/306,604

<151> 2001-07-19

<150> 60/315,412

<151> 2001-08-28

<160> 81

<170> PatentIn version 3.0

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 Pro Asp Ala Tyr Lys Ile Met Ser Gln Arg Cys Asn Asn Arg Thr Gln
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 195 200 205
 Thr Gly Phe Val Val Tyr Asp Gly Ala Leu Phe Phe Asn Lys Glu Arg
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Val Lys His Ser Asp Ala Val His Asp Leu Leu Leu Asp Val Ile Thr
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930 935 940
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Gly Val Gln Leu Tyr Ile Met Leu Val Glu Val Phe Glu Ser Glu His
965 970 975
Ser Arg Arg Lys Tyr Phe Tyr Leu Val Gly Tyr Gly Met Pro Ala Leu
980 985 990
Ile Val Ala Val Ser Ala Ala Val Asp Tyr Arg Ser Tyr Gly Thr Asp
995 1000 1005
Lys Val Cys Trp Leu Arg Leu Asp Thr Tyr Phe Ile Trp Ser Phe
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 Ile Cys Ile Phe Thr Phe Cys Phe Phe Arg Gly Leu Gln Ser Asp Arg
 965 970 975
 Asn Thr Ile His Lys Asn Leu Cys Ile Ser Leu Phe Val Ala Glu Leu
 980 985 990
 Leu Phe Leu Ile Gly Ile Asn Arg Thr Asp Gln Pro Ile Ala Cys Ala
 995 1000 1005
 Val Phe Ala Ala Leu Leu His Phe Phe Phe Leu Ala Ala Phe Thr
 1010 1015 1020
 Trp Met Phe Leu Glu Gly Val Gln Leu Tyr Ile Met Leu Val Glu
 1025 1030 1035
 Val Phe Glu Ser Glu His Ser Arg Arg Lys Tyr Phe Tyr Leu Val
 1040 1045 1050
 Gly Tyr Gly Met Pro Ala Leu Ile Val Ala Val Ser Ala Ala Val
 1055 1060 1065
 Asp Tyr Arg Ser Tyr Gly Thr Asp Lys Val Cys Trp Leu Arg Leu
 1070 1075 1080
 Asp Thr Tyr Phe Ile Trp Ser Phe Ile Gly Pro Ala Thr Leu Ile
 1085 1090 1095

Ile Met	Leu Asn Val	Ile Phe	Leu Gly Ile Ala	Leu Tyr Lys Met
1100		1105		1110
Phe His	His Thr Ala Ile	Leu Lys Pro Glu Ser	Gly Cys Leu Asp	
1115		1120	1125	
Asn Ile	Lys Ser Trp Val	Ile Gly Ala Ile Ala	Leu Leu Cys Leu	
1130		1135	1140	
Leu Gly	Leu Thr Trp Ala	Phe Gly Leu Met Tyr	Ile Asn Glu Ser	
1145		1150	1155	
Thr Val	Ile Met Ala Tyr	Leu Phe Thr Ile Phe	Asn Ser Leu Gln	
1160		1165	1170	
Gly Met	Phe Ile Phe Ile	Phe His Cys Val Leu	Gln Lys Lys Val	
1175		1180	1185	
Arg Lys	Glu Tyr Gly Lys	Cys Leu Arg Thr His	Cys Cys Ser Gly	
1190		1195	1200	
Lys Ser	Thr Glu Ser Ser	Ile Gly Ser Gly Lys	Thr Ser Gly Ser	
1205		1210	1215	
Arg Thr	Pro Gly Arg Tyr	Ser Thr Gly Ser Gln	Ser Arg Ile Arg	
1220		1225	1230	
Arg Met	Trp Asn Asp Thr	Val Arg Lys Gln Ser	Glu Ser Ser Phe	
1235		1240	1245	
Ile Thr	Gly Asp Ile Asn	Ser Ser Ala Ser Leu	Asn Arg Glu Gly	
1250		1255	1260	
Leu Leu	Asn Asn Ala Arg	Asp Thr Ser Val Met	Asp Thr Leu Pro	
1265		1270	1275	
Leu Asn	Gly Asn His Gly	Asn Ser Tyr Ser Ile	Ala Gly Gly Glu	
1280		1285	1290	
Tyr Leu	Ser Asn Cys Val	Gln Ile Ile Asp Arg	Gly Tyr Asn His	
1295		1300	1305	
Asn Glu	Thr Ala Leu Glu	Lys Lys Ile Leu Lys	Glu Leu Thr Ser	
1310		1315	1320	
Asn Tyr	Ile Pro Ser Tyr	Leu Asn Asn His Glu	Arg Ser Ser Glu	
1325		1330	1335	
Gln Asn	Arg Asn Met Met	Asn Lys Leu Val Asp	Asn Leu Gly Ser	
1340		1345	1350	
Gly Ser	Glu Asp Asp Ala	Ile Val Leu Asp Asp	Ala Ala Ser Phe	
1355		1360	1365	
Asn His	Glu Glu Ser Leu	Gly Leu Glu Leu Ile	His Glu Glu Ser	
1370		1375	1380	

Asp Ala Pro Leu Leu Pro Pro Arg Val Tyr Ser Thr Asp Asn His
 1385 1390 1395
 Gln Pro His His Tyr Ser Arg Arg Arg Leu Pro Gln Asp His Ser
 1400 1405 1410
 Glu Ser Phe Phe Pro Leu Leu Thr Asp Glu His Thr Glu Asp Pro
 1415 1420 1425
 Gln Ser Pro His Arg Asp Ser Leu Tyr Thr Ser Met Pro Ala Leu
 1430 1435 1440
 Ala Gly Val Pro Ala Ala Asp Ser Val Thr Thr Ser Thr Gln Thr
 1445 1450 1455
 Glu Ala Ala Ala Ala Lys Gly Gly Asp Ala Glu Asp Val Tyr Tyr
 1460 1465 1470
 Lys Ser Met Pro Asn Leu Gly Ser Arg Asn His Val His Pro Leu
 1475 1480 1485
 His Ala Tyr Tyr Gln Leu Gly Arg Gly Ser Ser Asp Gly Phe Ile
 1490 1495 1500
 Val Pro Pro Asn Lys Asp Gly Ala Ser Pro Glu Gly Thr Ser Lys
 1505 1510 1515
 Gly Pro Ala His Leu Val Thr Ser Leu
 1520 1525
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 Pro Pro Asn Leu Leu Glu Asn Leu Ser Pro Glu Asp Ser Val Leu Val
 20 25 30
 Arg Arg Ala Gln Phe Thr Phe Phe Asn Lys Thr Gly Leu Phe Gln Asp
 35 40 45
 Val Gly Pro Gln Arg Lys Thr Leu Val Ser Tyr Val Met Ala Cys Ser
 50 55 60
 Ile Gly Asn Ile Thr Ile Gln Asn Leu Lys Asp Pro Val Gln Ile Lys
 65 70 75 80
 Ile Lys His Thr Arg Thr Gln Glu Val His His Pro Ile Cys Ala Phe
 85 90 95
 Trp Asp Leu Asn Lys Asn Lys Ser Phe Gly Gly Trp Asn Thr Ser Gly

100	105	110
Cys Val Ala His Arg Asp Ser Asp Ala Ser Glu Thr Val Cys Leu Cys 115 120 125		
Asn His Phe Thr His Phe Gly Val Leu Met Asp Leu Pro Arg Ser Ala 130 135 140		
Ser Gln Leu Asp Ala Arg Asn Thr Lys Val Leu Thr Phe Ile Ser Tyr 145 150 155 160		
Ile Gly Cys Gly Ile Ser Ala Ile Phe Ser Ala Ala Thr Leu Leu Thr 165 170 175		
Tyr Val Ala Phe Glu Lys Leu Arg Arg Asp Tyr Pro Ser Lys Ile Leu 180 185 190		
Met Asn Leu Ser Thr Ala Leu Leu Phe Leu Asn Leu Leu Phe Leu Leu 195 200 205		
Asp Gly Trp Ile Thr Ser Phe Asn Val Asp Gly Leu Cys Ile Ala Val 210 215 220		
Ala Val Leu Leu His Phe Phe Leu Leu Ala Thr Phe Thr Trp Met Gly 225 230 235 240		
Leu Glu Ala Ile His Met Tyr Ile Ala Leu Val Lys Val Phe Asn Thr 245 250 255		
Tyr Ile Arg Arg Tyr Ile Leu Lys Phe Cys Ile Ile Gly Trp Gly Leu 260 265 270		
Pro Ala Leu Val Val Ser Val Val Leu Ala Ser Arg Asn Asn Asn Glu 275 280 285		
Val Tyr Gly Lys Glu Ser Tyr Gly Lys Glu Lys Gly Asp Glu Phe Cys 290 295 300		
Trp Ile Gln Asp Pro Val Ile Phe Tyr Val Thr Cys Ala Gly Tyr Phe 305 310 315 320		
Gly Val Met Phe Phe Leu Asn Ile Ala Met Phe Ile Val Val Met Val 325 330 335		
Gln Ile Cys Gly Arg Asn Gly Lys Arg Ser Asn Arg Thr Leu Arg Glu 340 345 350		
Glu Val Leu Arg Asn Leu Arg Ser Val Val Ser Leu Thr Phe Leu Leu 355 360 365		
Gly Met Thr Trp Gly Phe Ala Phe Phe Ala Trp Gly Pro Leu Asn Ile 370 375 380		
Pro Phe Met Tyr Leu Phe Ser Ile Phe Asn Ser Leu Gln Gly Leu Phe 385 390 395 400		
Ile Phe Ile Phe His Cys Ala Met Lys Glu Asn Val Gln Lys Gln Trp		

405										410					415						
Arg	Gln	His	Leu	Cys	Cys	Gly	Arg	Phe	Arg	Leu	Ala	Asp	Asn	Ser	Asp						
			420					425					430								
Trp	Ser	Lys	Thr	Ala	Thr	Asn	Ile	Ile	Lys	Lys	Ser	Ser	Asp	Asn	Leu						
		435					440					445									
Gly	Lys	Ser	Leu	Ser	Ser	Ser	Ser	Ile	Gly	Ser	Asn	Ser	Thr	Tyr	Leu						
	450					455					460										
Thr	Ser	Lys	Ser	Lys	Ser	Ser	Ser	Thr	Thr	Tyr	Phe	Lys	Arg	Asn	Ser						
465					470					475					480						
His	Thr	Asp	Ser	Ala	Ser	Met	Asp	Lys	Ser	Leu	Ser	Lys	Leu	Ala	His						
				485					490					495							
Ala	Asp	Gly	Asp	Gln	Thr	Ser	Ile	Ile	Pro	Val	His	Gln	Val	Ile	Asp						
			500					505					510								
Lys	Val	Lys	Gly	Tyr	Cys	Asn	Ala	His	Ser	Asp	Asn	Phe	Tyr	Lys	Asn						
		515					520					525									
Ile	Ile	Met	Ser	Asp	Thr	Phe	Ser	His	Ser	Thr	Lys	Phe									
	530					535					540										

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 <213> Caenorhabditis elegans

<400> 11

Met	Ala	Thr	Ala	Ser	Thr	Glu	Ile	Ser	Glu	Phe	Ser	Glu	Ala	Ile	Glu						
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Ser	Thr	Phe	Asp	Leu	Asp	Phe	Thr	Ala	His	Gln	Thr	Glu	Ile	Ile	Gly						
			20					25					30								
Thr	Tyr	Trp	Asn	Leu	Arg	Ala	Leu	Leu	Arg	Leu	His	Arg	Ser	Leu	Val						
		35					40					45									
Ala	Ile	Asp	His	Val	Ser	Gln	Lys	Ser	Phe	Trp	Glu	Arg	Tyr	Asn	His						
	50					55					60										
Trp	Ile	Gln	Leu	Ser	Met	Leu	Val	Ser	Asn	Gln	Asn	Val	Asn	Leu	Cys						
65					70					75				80							
Gln	Ser	Asn	Ile	Cys	Gln	Asn	Gly	Gly	Thr	Cys	Leu	Val	Ala	Ser	Ser						
				85				90						95							
Val	Pro	Ala	Thr	Ala	Thr	Cys	Pro	Lys	Asn	Ser	Ile	Tyr	Tyr	Met	Gly						
		100						105					110								
Ser	Cys	Tyr	Val	Phe	Asp	Thr	Thr	Leu	Arg	Asn	Trp	Asn	Asp	Ala	Ala						
		115					120					125									

Leu Tyr Cys Asn Asn Met Asn Ser Ala Thr Leu Pro Leu Val Glu Ser
 130 135 140
 Ala Glu Asp Gln Ala Phe Phe Ala Gly Tyr Leu Gln Ala Met Ile Pro
 145 150 155 160
 Ser Asn Pro Pro Ala Asp Met Arg Pro Pro Pro Asp Gly Ile Trp Thr
 165 170 175
 Ala Val Arg Gly Val Asn Asn Val Thr Arg Ala Ser Trp Val Tyr Tyr
 180 185 190
 Pro Gly Ser Phe Leu Val Thr Asp Thr Phe Trp Ala Pro Gln Glu Pro
 195 200 205
 Asn Ile Tyr Val Asn Tyr Asn Asp Val Cys Val Ala Leu Gln Ser Asp
 210 215 220
 Ser Phe Tyr Arg Glu Trp Thr Thr Ala Leu Cys Thr Ile Leu Lys Tyr
 225 230 235 240
 Thr Val Cys Lys Val Ala Pro Thr Gln Ile Gln Ala Lys Tyr Val Ala
 245 250 255
 Gln Cys Ser Cys Pro Asn Gly Tyr Gly Gly Gln Thr Cys Glu Thr Gln
 260 265 270
 Ser Thr Thr Asn Gln Gln Ala Ser Thr Gln Arg Thr Cys Gly Ser Asn
 275 280 285
 Asp Phe Gln Phe Ser Cys Pro Asn Asp Gln Thr Ile Thr Val Asp Phe
 290 295 300
 Ala Ser Phe Gly Ala Gln Gly Gly Ser Ile Ile Thr Ser Pro Pro Asp
 305 310 315 320
 Ala Leu Leu Gln Gln Ile Val Gln Lys Val Asn Ala Glu Thr Lys Lys
 325 330 335
 Thr Val Asn Phe Trp Ile Gly Thr Pro Asn Asn Cys Gln Leu Leu Met
 340 345 350
 Val Thr Gly Ser Ser Thr Ser Tyr Ser Gln Cys Pro Ser Ser Pro Ser
 355 360 365
 Ser Thr Ala Asn Val Ile Cys Ser Thr Val Pro Gln Ser Thr Ala Ser
 370 375 380
 Val Ser Ala Arg Pro Thr Gln Ser Ala Pro Val Asp Pro Val Ser Gln
 385 390 395 400
 Thr Met Ala Arg Arg Glu Val Tyr Thr Gly Val Gln Pro Ile Ala Ser
 405 410 415
 Ala Leu Gly Gly Gln Ser Lys Lys Thr Asn Arg Lys Leu Asn Asn Ile
 420 425 430

Cys Gln Thr Lys Ile Gly Ala Pro Leu Ser Leu Phe Leu Phe Ser Arg
 435 440 445
 Asn Glu Val Ile Thr Gly Phe Val Cys Ile Ser Leu Ile Ser Ala Ser
 450 455 460
 Pro Gln Ile Ile Tyr Tyr Leu Cys Ala Val Ser Leu Ile Cys His Pro
 465 470 475 480
 Ser Val Pro Asp Ser Ile Asn Lys Pro Arg Tyr Cys Lys Lys Glu Lys
 485 490 495
 Lys Asp Gly Ile Thr Tyr Glu Gln Thr Arg Ala Cys Met Leu His Glu
 500 505 510
 Gln Pro Cys Pro Asp Pro Gln Asn Val Glu Gly Thr Val Thr Arg Tyr
 515 520 525
 Cys Asn Cys Gln Thr Ala Lys Trp Glu Thr Pro Asp Thr Thr Asn Cys
 530 535 540
 Thr His Arg Trp Val Ala Glu Met Glu Thr Ala Ile Lys Asp Asn Gln
 545 550 555 560
 Pro Val Glu Asp Ile Ser Ser Thr Val Asn Arg Gln Leu Lys Ser Thr
 565 570 575
 Ile Glu Arg Thr Leu Phe Gly Gly Asp Ile Thr Gly Thr Val Arg Leu
 580 585 590
 Ser Asn Asp Met Leu Ser Leu Ala Arg Asn Gln Phe Ser Val Leu Asn
 595 600 605
 Asp Arg Asn Leu Arg Glu Asn Lys Ala Arg Asn Phe Thr Glu Asn Leu
 610 615 620
 Gly Gly Ser Gly Asp Gln Leu Leu Ser Pro Val Ala Ala Thr Val Trp
 625 630 635 640
 Asp Gln Leu Ser Ser Thr Ile Arg Ile Gln His Ala Ser Lys Leu Met
 645 650 655
 Ser Val Leu Glu Gln Ser Val Leu Leu Leu Gly Asp Tyr Met Thr Asp
 660 665 670
 Gln Lys Leu Asn Leu Gln Tyr Ile Asn Trp Ala Met Glu Val Glu Arg
 675 680 685
 Ser Glu Pro Glu Val Gln Thr Phe Gly Ala Ala Ala Ser Pro Asn Val
 690 695 700
 Gln Asp Asp Met Gly Met Met Arg Val Met Ala Ala Ala Pro Pro Ala
 705 710 715 720
 Pro Gln Pro Glu Thr Asn Thr Thr Ile Met Phe Pro Ser Leu Lys Leu
 725 730 735

Ser Pro Thr Ile Thr Leu Pro Ser Ala Ser Leu Leu Ser Ser Leu Ala
 740 745 750
 Ser Pro Thr Pro Val Ala Gly Gly Gly Pro Ser Ile Leu Ser Ser Phe
 755 760 765
 Gln Asp Asp Thr Pro Val Gly Met Ala Ser Thr Pro Asn Leu Asn Arg
 770 775 780
 Asn Pro Val Lys Leu Gly Tyr Tyr Ala Phe Ala Gly Phe Gly Gln Leu
 785 790 795 800
 Leu Asn Asn Asn Asn Asp His Thr Leu Ile Asn Ser Gln Val Ile Gly
 805 810 815
 Ala Ser Ile Gln Asn Ala Thr Gln Ser Val Thr Leu Pro Val Asp His
 820 825 830
 Pro Val Thr Phe Thr Phe Gln His Leu Thr Thr Lys Gly Val Ser Asn
 835 840 845
 Pro Arg Cys Val Tyr Trp Asp Leu Met Glu Ser Lys Trp Ser Thr Leu
 850 855 860
 Gly Cys Thr Leu Ile Ala Thr Ser Ser Asn Ser Ser Gln Cys Ser Cys
 865 870 875 880
 Thr His Leu Thr Ser Phe Ala Ile Leu Met Asp Ile Ser Gly Gln Val
 885 890 895
 Gly Arg Leu Ser Gly Gly Leu Ala Ser Ala Leu Asp Val Val Ser Thr
 900 905 910
 Ile Gly Cys Ala Ile Ser Ile Val Cys Leu Ala Leu Ser Val Cys Val
 915 920 925
 Phe Thr Phe Phe Arg Asn Leu Gln Asn Val Arg Asn Ser Ile His Arg
 930 935 940
 Asn Leu Cys Leu Cys Leu Leu Ile Ala Glu Leu Val Phe Val Ile Gly
 945 950 955 960
 Met Asp Arg Thr Gly Asn Arg Thr Gly Cys Gly Val Val Ala Ile Leu
 965 970 975
 Leu His Tyr Phe Phe Leu Ser Ser Phe Cys Trp Met Leu Leu Glu Gly
 980 985 990
 Tyr Gln Leu Tyr Met Met Leu Ile Gln Val Phe Glu Pro Asn Arg Thr
 995 1000 1005
 Arg Ile Phe Leu Tyr Tyr Leu Phe Cys Tyr Gly Thr Pro Ala Val
 1010 1015 1020
 Val Val Ala Ile Ser Ala Gly Ile Lys Trp Glu Asp Tyr Gly Thr
 1025 1030 1035

Asp	Ser	Tyr	Cys	Trp	Ile	Asp	Thr	Ser	Thr	Pro	Thr	Ile	Trp	Ala
1040						1045					1050			
Phe	Val	Ala	Pro	Ile	Ile	Val	Ile	Ile	Ala	Ala	Asn	Ile	Ile	Phe
1055						1060					1065			
Leu	Leu	Ile	Ala	Leu	Lys	Val	Val	Leu	Ser	Val	Gln	Ser	Arg	Asp
1070						1075					1080			
Arg	Thr	Lys	Trp	Gly	Arg	Ile	Ile	Gly	Trp	Leu	Lys	Gly	Ser	Ala
1085						1090					1095			
Thr	Leu	Leu	Cys	Leu	Leu	Gly	Ile	Thr	Trp	Ile	Phe	Gly	Phe	Leu
1100						1105					1110			
Thr	Ala	Val	Lys	Gly	Gly	Thr	Gly	Thr	Ala	Phe	Ala	Trp	Ile	Phe
1115						1120					1125			
Thr	Ile	Leu	Asn	Cys	Thr	Gln	Gly	Ile	Phe	Ile	Phe	Val	Leu	His
1130						1135					1140			
Val	Val	Leu	Asn	Glu	Lys	Val	Arg	Ala	Ser	Ile	Val	Arg	Trp	Leu
1145						1150					1155			
Arg	Thr	Gly	Ile	Cys	Cys	Leu	Pro	Glu	Thr	Ser	Ser	Ala	Ala	Tyr
1160						1165					1170			
Asn	Ser	Arg	Ser	Phe	Leu	Ser	Ser	Arg	Gln	Arg	Ile	Leu	Asn	Met
1175						1180					1185			
Ile	Lys	Val	Asn	Gly	His	Ser	Tyr	Pro	Ser	Thr	Ala	Ser	Thr	Asp
1190						1195					1200			
Asp	Lys	Glu	Lys	Gln	Leu	Thr	Pro	Ile	Thr	Lys	Thr	Thr	Asp	Trp
1205						1210					1215			
Leu	Ser	Arg	Leu	Pro	Asn	Gln	Asp	Ser	Val	Ser	Ile	Pro	Glu	Ser
1220						1225					1230			
Asn	Phe	Asn	Asn	Leu	Asn	Gly	Thr	Leu	Glu	Asn	Ser	Asn	Leu	Asn
1235						1240					1245			
Ser	Ala	Glu	Ile	Lys	Glu	Glu	Asp	Glu	Ile	Pro	Glu	Leu	Arg	Arg
1250						1255					1260			
Arg	Val	Thr	Val	Asp	Leu	Asn	Pro	Met	Ile	Val	Ser	Asn	Asn	Glu
1265						1270					1275			
Ile	Glu	Arg	Met	Ser	His	Ala	Ser	Ser	Asp	Pro	Arg	Gly	Ser	Gln
1280						1285					1290			
Ile	Ile	Glu	Val	Thr	Ala	Val	Glu	Lys	Lys	Ala	Pro	Val	Lys	Arg
1295						1300					1305			
Ile	Lys	Phe	Pro	Leu	Gly	Ala	Lys	Gln	Ser	Glu	Arg	Gly	Ser	Gln
1310						1315					1320			

His	Arg	Thr	Lys	Ala	Lys	His	Gly	Thr	Gly	Thr	Leu	Val	Ser	Pro	1325	1330	1335
Trp	His	Ile	Val	Thr	Ala	Ala	His	Leu	Ile	Gly	Ile	Ser	Glu	Asp	1340	1345	1350
Pro	Leu	Pro	Asp	Cys	Asp	Thr	Gly	Asn	Leu	Arg	Glu	Ala	Tyr	Phe	1355	1360	1365
Val	Arg	Asp	Tyr	Lys	Asn	Phe	Val	Ala	Phe	Val	Asn	Val	Thr	Cys	1370	1375	1380
Ala	Val	Pro	Glu	Met	Cys	Lys	Gly	Leu	His	Arg	Lys	Asp	Met	Phe	1385	1390	1395
Lys	Pro	Leu	Ala	Ile	Lys	Ser	Leu	Tyr	Ile	Arg	Lys	Gly	Tyr	Val	1400	1405	1410
Gly	Asp	Gly	Cys	Ile	Asp	Arg	Glu	Ser	Phe	Asn	Asp	Ile	Ala	Val	1415	1420	1425
Phe	Glu	Leu	Glu	Glu	Pro	Ile	Glu	Phe	Ser	Lys	Asp	Ile	Phe	Pro	1430	1435	1440
Ala	Cys	Leu	Pro	Ser	Ala	Pro	Lys	Ile	Pro	Arg	Ile	Arg	Glu	Thr	1445	1450	1455
Gly	Tyr	Lys	Leu	Phe	Gly	Tyr	Gly	Arg	Asp	Pro	Ser	Asp	Ser	Val	1460	1465	1470
Leu	Glu	Ser	Gly	Lys	Leu	Lys	Ser	Leu	Tyr	Ser	Phe	Val	Ala	Glu	1475	1480	1485
Cys	Ser	Asp	Asp	Phe	Pro	Tyr	Gly	Gly	Val	Tyr	Cys	Thr	Ser	Ala	1490	1495	1500
Val	Asn	Arg	Gly	Leu	Ser	Cys	Asp	Gly	Asp	Ser	Gly	Ser	Gly	Val	1505	1510	1515
Val	Arg	Thr	Ser	Asp	Thr	Arg	Asn	Val	Gln	Val	Leu	Val	Gly	Val	1520	1525	1530
Leu	Ser	Ala	Gly	Met	Pro	Cys	Pro	Glu	Leu	Tyr	Asp	Thr	His	Asn	1535	1540	1545
Arg	Gln	Arg	Gln	Gln	Arg	Arg	Gln	Leu	Thr	Gln	Glu	Thr	Asp	Leu	1550	1555	1560
Leu	Val	Asp	Val	Ser	Ala	His	Val	Asp	Phe	Phe	Cys	Thr	Cys	Cys	1565	1570	1575
Gly	Met	Cys	Ser												1580		
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<213> Homo sapiens

<400> 12

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Asn Ile Ala Ile Gln Ser Ala Asn Phe Ser Ser Glu Asn Ala Val Gly
20 25 30
Pro Ser Asn Val Arg Phe Ser Val Gln Lys Gly Ala Ser Ser Ser Leu
35 40 45
Val Ser Ser Ser Thr Phe Ile His Thr Asn Val Asp Gly Leu Asn Pro
50 55 60
Asp Ala Gln Thr Glu Leu Gln Val Leu Leu Asn Met Thr Lys Asn Tyr
65 70 75 80
Thr Lys Thr Cys Gly Phe Val Val Tyr Gln Asn Asp Lys Leu Phe Gln
85 90 95
Ser Lys Thr Phe Thr Ala Lys Ser Asp Phe Ser Gln Lys Ile Ile Ser
100 105 110
Ser Lys Thr Asp Glu Asn Glu Gln Asp Gln Ser Ala Ser Val Asp Met
115 120 125
Val Phe Ser Pro Lys Tyr Asn Gln Lys Glu Phe Gln Leu Tyr Ser Tyr
130 135 140
Ala Cys Val Tyr Trp Asn Leu Ser Ala Lys Asp Trp Asp Thr Tyr Gly
145 150 155 160
Cys Gln Lys Asp Lys Gly Thr Asp Gly Phe Leu Arg Cys Arg Cys Asn
165 170 175
His Thr Thr Asn Phe Ala Val Leu Met Thr Phe Lys Lys Asp Tyr Gln
180 185 190
Tyr Pro Lys Ser Leu Asp
195

<210> 13

<211> 10

<212> PRT

<213> Homo sapiens

<400> 13

Gln Ile Val Thr Arg Lys Val Arg Lys Thr
1 5 10

<210> 14

<211> 38

<212> PRT

<213> Homo sapiens

<400> 14

Glu Asn Ser Asn Lys Asn Leu Gln Thr Ser Asp Gly Asp Ile Asn Asn
1 5 10 15

Ile Asp Phe Asp Asn Asn Asp Ile Pro Arg Thr Asp Thr Ile Asn Ile
20 25 30

Pro Asn Pro Met Cys Thr
35

<210> 15

<211> 10

<212> PRT

<213> Homo sapiens

<400> 15

Ile Arg Thr Met Lys Pro Leu Pro Arg His
1 5 10

<210> 16

<211> 41

<212> PRT

<213> Homo sapiens

<400> 16

Thr Val Gly Val Ile Tyr Ser Gln Asn Gly Asn Asn Pro Gln Trp Glu
1 5 10 15

Leu Asp Tyr Arg Gln Glu Lys Ile Cys Trp Leu Ala Ile Pro Glu Pro
20 25 30

Asn Gly Val Ile Lys Ser Pro Leu Leu
35 40

<210> 17

<211> 25

<212> PRT

<213> Homo sapiens

<400> 17

Thr Ile Ser Ile Lys Val Leu Trp Lys Asn Asn Gln Asn Leu Thr Ser
1 5 10 15

Thr Lys Lys Val Ser Ser Met Lys Lys
20 25

<210> 18

<211> 6

<212> PRT

<213> Homo sapiens

<400> 18

Asn Asp Asp Ser Ile Arg
1 5

<210> 19
<211> 78
<212> PRT
<213> Homo sapiens

<400> 19

Tyr Thr Val Arg Thr Lys Val Phe Gln Ser Glu Ala Ser Lys Val Leu
1 5 10 15

Met Leu Leu Ser Ser Ile Gly Arg Arg Lys Ser Leu Pro Ser Val Thr
20 25 30

Arg Pro Arg Leu Arg Val Lys Met Tyr Asn Phe Leu Arg Ser Leu Pro
35 40 45

Thr Leu His Glu Arg Phe Arg Leu Leu Glu Thr Ser Pro Ser Thr Glu
50 55 60

Glu Ile Thr Leu Ser Glu Ser Asp Asn Ala Lys Glu Ser Ile
65 70 75

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<211> 38
<212> DNA
<213> Artificial

<220>
<223> HGPRBMY6 5' PRIMER

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38

<210> 21
<211> 66
<212> DNA
<213> Artificial

<220>
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actttc

60

66

<210> 22
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<220>
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 <400> 22
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23

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22

<210> 24
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 <220>
 <223> GAPDH-F3 forward primer

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17

<210> 25
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 <212> DNA
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19

<210> 26
 <211> 28
 <212> DNA
 <213> Homo sapiens

 <400> 26
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28

<210> 27
 <211> 13
 <212> PRT
 <213> Homo sapiens

<400> 27

Gln Ser Lys Thr Phe Thr Ala Lys Ser Asp Phe Ser Gln
1 5 10

<210> 28

<211> 13

<212> PRT

<213> Homo sapiens

<400> 28

Ala Lys Ser Asp Phe Ser Gln Lys Ile Ile Ser Ser Lys
1 5 10

<210> 29

<211> 13

<212> PRT

<213> Homo sapiens

<400> 29

Ser Gln Lys Ile Ile Ser Ser Lys Thr Asp Glu Asn Glu
1 5 10

<210> 30

<211> 13

<212> PRT

<213> Homo sapiens

<400> 30

Val Asp Met Val Phe Ser Pro Lys Tyr Asn Gln Lys Glu
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Leu Glu Trp Gly Ser Asp Val Phe Tyr Asp Val Tyr Asp Cys Cys
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Arg Ile Asp Ser Cys Ala Lys Tyr Phe Leu Arg Ser Cys Asp
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Cys Leu Arg Ser Gly Thr Gly Cys Ala Phe Gln Leu Tyr Arg Phe
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Phe Arg Val Ser Arg Val Trp Asn Pro Pro Ser Phe Asp Ser Ala
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37